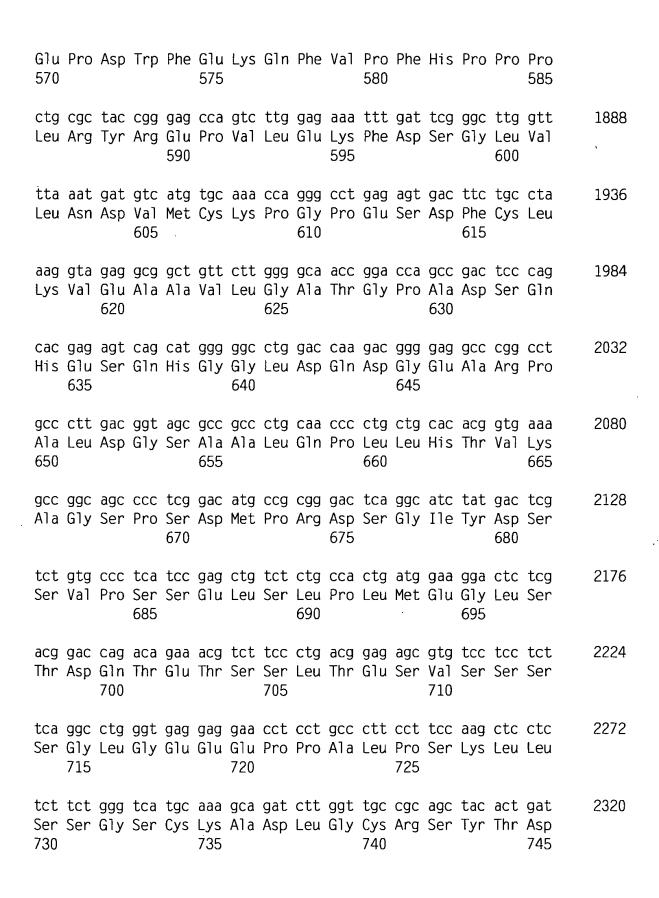
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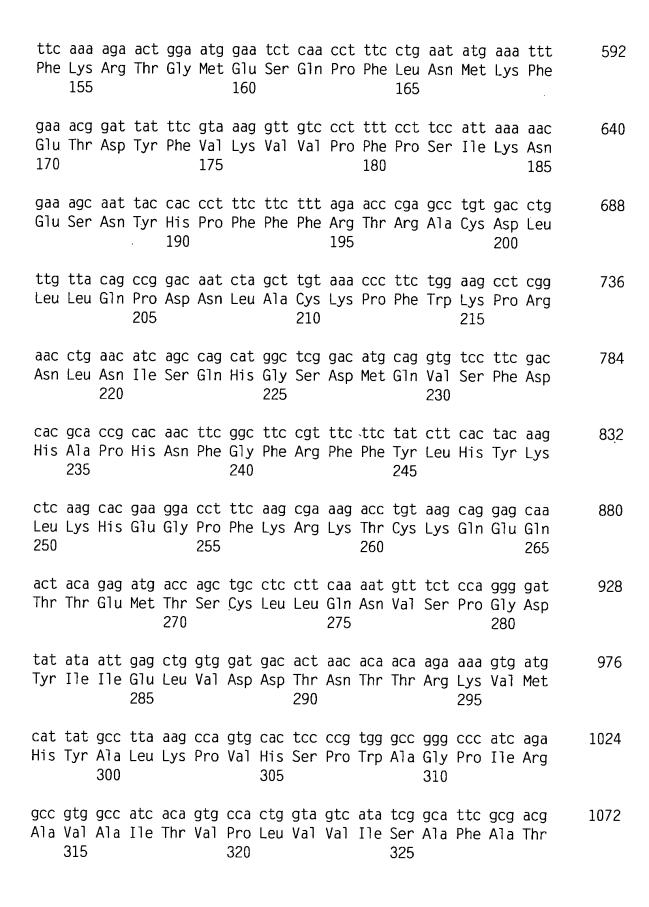
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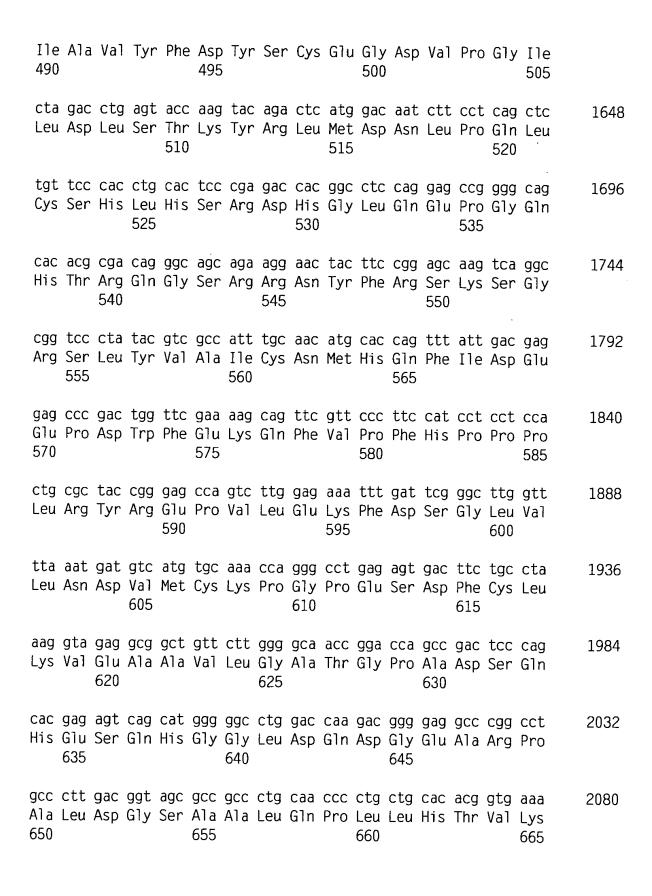
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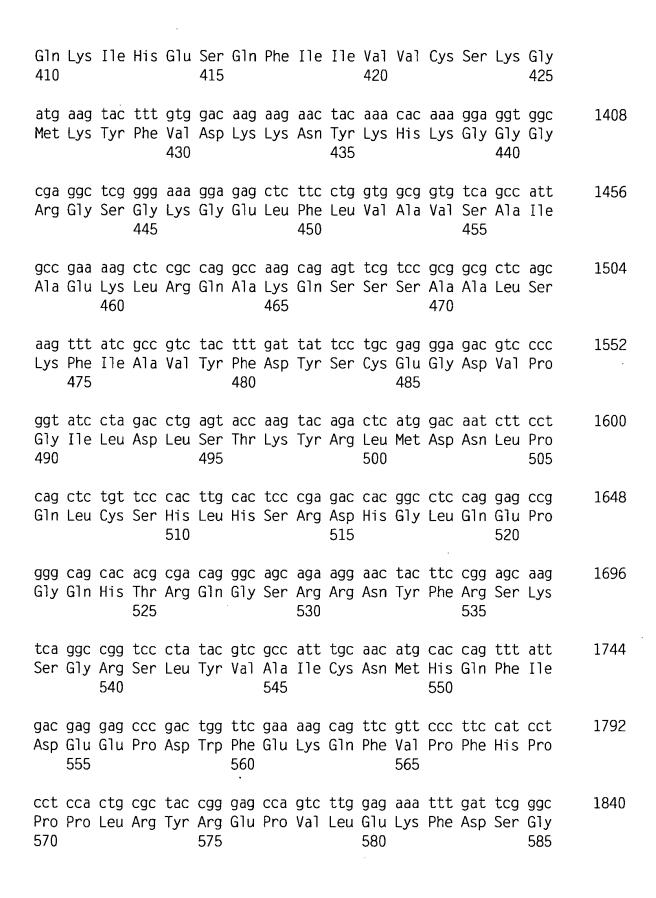
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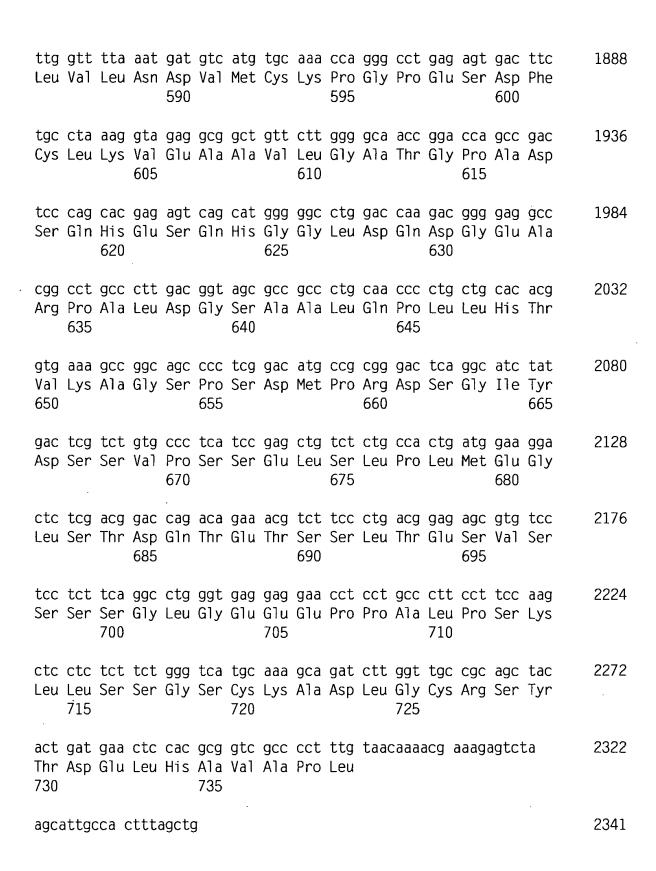
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				act Thr		-	-	-	_	-			-			1072
				gat Asp												1120
				gag Glu 350				-		_	_	_			_	1168
			Lys	gat Asp	Gly	Gln	Asn	His	_	Asn	Val	-	_	Cys		1216
				cag Gln												1264
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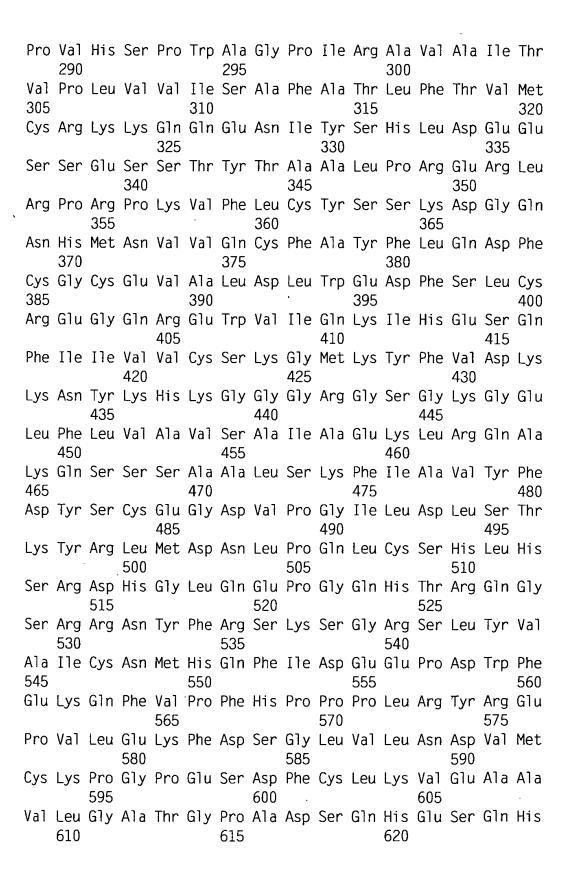




<210> 8 <211> 739 <212> PRT <213> Homo sapiens

<400> 8

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Gly Gly Leu Asp Gln Asp Gly Glu Ala Arg Pro Ala Leu Asp Gly Ser 625 630 640 635 Ala Ala Leu Gln Pro Leu Leu His Thr Val Lys Ala Gly Ser Pro Ser 645 650 Asp Met Pro Arg Asp Ser Gly Ile Tyr Asp Ser Ser Val Pro Ser Ser 660 665 670 Glu Leu Ser Leu Pro Leu Met Glu Gly Leu Ser Thr Asp Gln Thr Glu 675 680 685 Thr Ser Ser Leu Thr Glu Ser Val Ser Ser Ser Gly Leu Gly Glu 695 700 Glu Glu Pro Pro Ala Leu Pro Ser Lys Leu Leu Ser Ser Gly Ser Cys 705 710 715 Lys Ala Asp Leu Gly Cys Arg Ser Tyr Thr Asp Glu Leu His Ala Val 725 730 735 Ala Pro Leu <210> 9 <211> 2217 <212> DNA <213> Artificial Sequence <220> <223> This degenerate nucleotide sequence encodes the amino acid sequence of SEQ ID NO:8. <221> misc feature <222> (1)...(2217) <223> n = A,T,C or G<400> 9 atggcnccnt ggytncaryt ntgywsngtn ttyttyacng tnaaygcntg yytnaayggn wsncarytng cngtngcngc nggnggnwsn ggnmgngcnm gngqngcnga yacntqyggn 120 tggmgnggng tnggnccngc nwsnmgnaay wsnggnytnt ayaayathac nttyaartay 180 gayaaytgya cnacntayyt naaycengtn ggnaarcayg tnathgenga ygenearaay 240 athacnathw sncartaygc ntgycaygay cargtngcng tnacnathyt ntggwsnccn 300 ggngcnytng gnathgartt yytnaarggn ttymgngtna thytngarga rytnaarwsn 360 garggnmgnc artgycarca rytnathytn aargayccna arcarytnaa ywsnwsntty 420

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gtnaargtng tnccnttycc nwsnathaar aaygarwsna aytaycaycc nttyttytty

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gcnccncaya ayttyggntt ymgnttytty tayytncayt ayaarytnaa rcaygarggn

centtyaarm gnaaraentg yaareargar caraenaeng araenaenws ntgyytnytn

60

480

540

600

660

720



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<220>

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<212> DNA

<213> Mouse

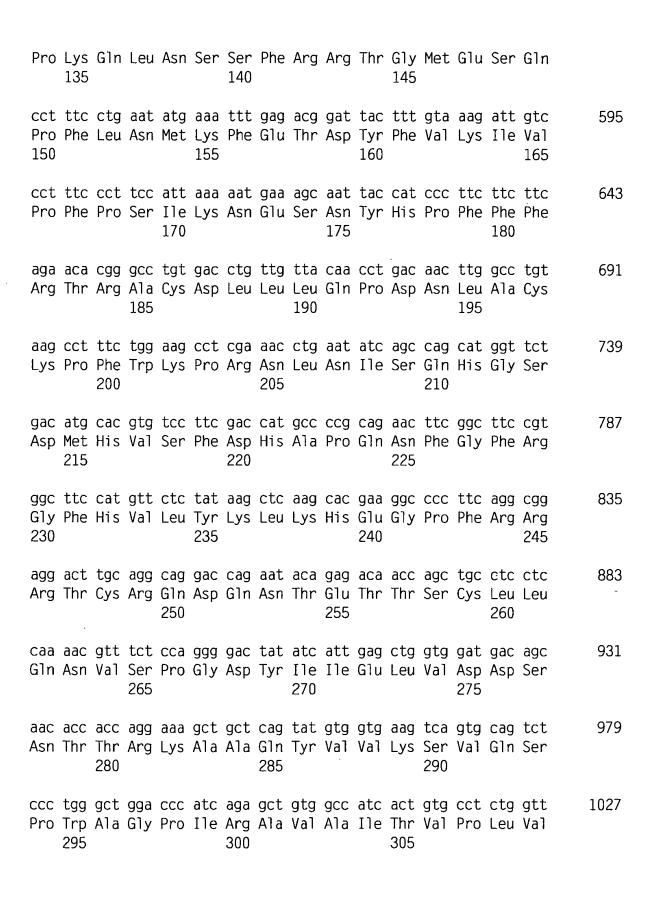
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	_			_						cgc Arg					-	211
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999 Gly											_	-		_		451
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_	gtc Val			_					_		_			_		1219
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	gct Ala	_	_	-		-	-		_		-	-			-	1315
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	aaa Lys														-	1459
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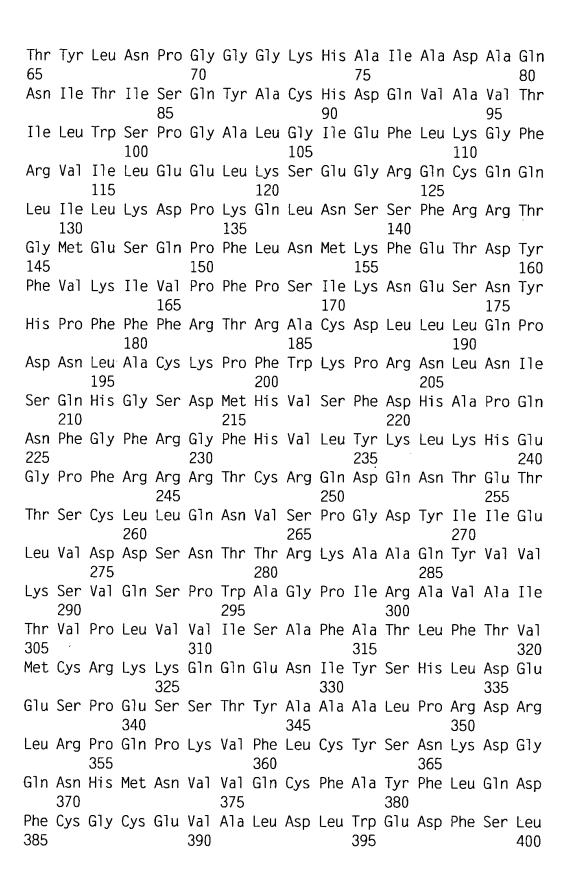


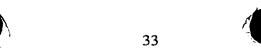
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acc gag agt gta tct tcc tcc tct ggc cta ggt Thr Glu Ser Val Ser Ser Ser Ser Gly Leu Gly 695 700		2227
acc ctc cct tcc aag ctc ttt gcc tct ggg gtg Thr Leu Pro Ser Lys Leu Phe Ala Ser Gly Val 710 715 720		2275
tgc cac agc cac act gac gaa ctg caa gcg ctt Cys His Ser His Thr Asp Glu Leu Gln Ala Leu 730 735		2317
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Ala Arg Gly Ala Asp Thr Cys Gly Trp Arg Gly 35 40	Val Gly Pro Ala Ser	
Arg Asn Ser Gly Leu His Asn Ile Thr Phe Arg	45 Tyr Asp Asn Cys Thr	





Cys Arg Glu Gly Gln Arg Glu Trp Ala Ile Gln Lys Ile His Glu Ser Gln Phe Ile Ile Val Val Cys Ser Lys Gly Met Lys Tyr Phe Val Asp 420 425 430 Lys Lys Asn Phe Arg His Lys Gly Gly Ser Arg Gly Glu Ala Gln Gly 440 445 Glu Phe Phe Leu Val Ala Val Ala Ala Ile Ala Glu Lys Leu Arg Gln 455 460 Ala Lys Gln Ser Ser Ser Ala Ala Leu Arg Lys Phe Ile Ala Val Tyr 465 470 475 Phe Asp Tyr Ser Cys Glu Gly Asp Val Pro Cys Ser Leu Asp Leu Ser 485 490 Thr Lys Tyr Lys Leu Met Asp His Leu Pro Glu Leu Cys Ala His Leu 500 505 510 His Ser Gly Glu Gln Glu Val Leu Gly Gln His Pro Gly His Ser Ser 515 520 525 Arg Arg Asn Tyr Phe Arg Ser Lys Ser Gly Arg Ser Leu Tyr Val Ala 535 Ile Cys Asn Met His Gln Phe Ile Asp Glu Glu Pro Asp Trp Phe Glu 550 555 560 Lys Gln Phe Ile Pro Phe Gln His Pro Pro Val Arg Tyr Gln Glu Pro 565 570 Val Leu Glu Lys Phe Asp Ser Gly Leu Val Leu Asn Asp Val Ile Ser 580 585 Lys Pro Gly Pro Glu Ser Asp Phe Cys Arg Lys Val Glu Ala Cys Val 595 600 605 Leu Gly Ala Ala Gly Pro Ala Asp Ser Tyr Ser Tyr Leu Glu Ser Gln 615 620 His Val Gly Leu Asp Gln Asp Thr Glu Ala Gln Pro Ser Cys Asp Ser 630 635 Ala Pro Ala Leu Gln Pro Leu Leu His Ala Val Lys Ala Gly Ser Pro 645 650 Ser Glu Met Pro Arg Asp Ser Gly Ile Tyr Asp Ser Ser Val Pro Ser 660 665 Ser Glu Leu Ser Leu Pro Leu Met Glu Gly Leu Ser Pro Asp Gln Ile 675 680 685 Glu Thr Ser Ser Leu Thr Glu Ser Val Ser Ser Ser Gly Leu Gly 690 695 700 Glu Glu Asp Pro Pro Thr Leu Pro Ser Lys Leu Phe Ala Ser Gly Val 710 715 Ser Arg Glu His Gly Cys His Ser His Thr Asp Glu Leu Gln Ala Leu 725 730 735





## Ala Pro Leu

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<211> 2217

<212> DNA

<213> Artificial Sequence

<220>

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<221> misc feature

<222> (1)...(2217)

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